SEQUENCE LISTING

<110> MOECKEL, BETTINA

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<141> 2001-08-02

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<151> 2000-09-02

<150> DE 10109690.9

<151> 2001-02-28

<150> US 60/294,252

<151> 2001-05-31

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376

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Phe Asp Ser Ala Glu His Ala Lys Gln Arg Phe Ala Leu Glu Asp Leu

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45

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| 80 85 90 | |
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| cta ttc ctt atc act ctt aac cgc ctg ggt atc gat gtt tcc ttc gtg Leu Phe Leu Ile Thr Leu Asn Arg Leu Gly Ile Asp Val Ser Phe Val 125 130 135 | 616 |
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| ctg atc atc gac aac acc atc gct acc gca gcg ctc gtg cgc ccg ctc Leu Ile Ile Asp Asn Thr Ile Ala Thr Ala Ala Leu Val Arg Pro Leu 190 195 200 | 808 |
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| _ | | | | | | | | | | | | | | | | | |
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| | | | | | | | | cgc Arg | | | | | | | | | 1144 |
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| | | | | | | | | tac Tyr | | | | | | | | | 1240 |
| | aag Lys | tac Tyr | acc Thr 350 | ggc Gly | tcc Ser | gtt Val | ctc Leu | acc Thr 355 | ttc Phe | gag Glu | atc Ile | aag Lys | ggc Gly 360 | ggc Gly | aag Lys | gat Asp | 1288 |
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| | tcc Ser | acc Thr | gtc Val | cgc Arg 415 | ctg Leu | tcc Ser | gtt Val | ggc Gly | atc Ile 420 | gag Glu | acc Thr | att Ile | gat Asp | gat Asp 425 | atc Ile | atc Ile | 1480 |
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His Ala Lys Gln Arg Phe Ala Leu Glu Asp Leu Gly Pro Val Tyr Ser 50 55 60

Arg Leu Thr Asn Pro Thr Val Glu Ala Leu Glu Asn Arg Ile Ala Ser 65 70 75 80

Leu Glu Gly Val His Ala Val Ala Phe Ser Ser Gly Gln Ala Ala 85 90 95

Thr Thr Asn Ala Ile Leu Asn Leu Ala Gly Ala Gly Asp His Ile Val 100 105 110

Thr Ser Pro Arg Leu Tyr Gly Gly Thr Glu Thr Leu Phe Leu Ile Thr 115 120 125

Leu Asn Arg Leu Gly Ile Asp Val Ser Phe Val Glu Asn Pro Asp Asp 130 135 140

Pro Glu Ser Trp Gln Ala Ala Val Gln Pro Asn Thr Lys Ala Phe Phe 145 150 155 160

Gly Glu Thr Phe Ala Asn Pro Gln Ala Asp Val Leu Asp Ile Pro Ala Val Ala Glu Val Ala His Arg Asn Ser Val Pro Leu Ile Ile Asp Asn Thr Ile Ala Thr Ala Ala Leu Val Arg Pro Leu Glu Leu Gly Ala Asp Val Val Ala Ser Leu Thr Lys Phe Tyr Thr Gly Asn Gly Ser Gly Leu Gly Gly Val Leu Ile Asp Gly Gly Lys Phe Asp Trp Thr Val Glu Lys Asp Gly Lys Pro Val Phe Pro Tyr Phe Val Thr Pro Asp Ala Ala Tyr His Gly Leu Lys Tyr Ala Asp Leu Gly Ala Pro Ala Phe Gly Leu Lys Val Arg Val Gly Leu Leu Arg Asp Thr Gly Ser Thr Leu Ser Ala Phe Asn Ala Trp Ala Ala Val Gln Gly Ile Asp Thr Leu Ser Leu Arg Leu Glu Arg His Asn Glu Asn Ala Ile Lys Val Ala Glu Phe Leu Asn Asn His Glu Lys Val Glu Lys Val Asn Phe Ala Gly Leu Lys Asp Ser Pro Trp Tyr Ala Thr Lys Glu Lys Leu Gly Leu Lys Tyr Thr Gly Ser Val Leu Thr Phe Glu Ile Lys Gly Gly Lys Asp Glu Ala Trp Ala Phe

Ile Asp Ala Leu Lys Leu His Ser Asn Leu Ala Asn Ile Gly Asp Val 370 380

Arg Ser Leu Val Val His Pro Ala Thr Thr Thr His Ser Gln Ser Asp 385 390 395 400

Glu Ala Gly Leu Ala Arg Ala Gly Val Thr Gln Ser Thr Val Arg Leu 405 410 415

Ser Val Gly Ile Glu Thr Ile Asp Asp Ile Ile Ala Asp Leu Glu Gly 420 425 430

Gly Phe Ala Ala Ile 435